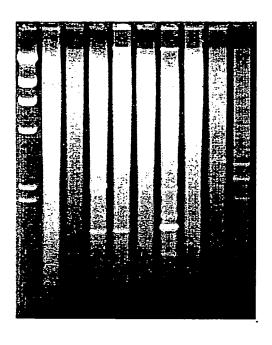
Fig. 1

Fig. A

Fig. B

1 2 3 4 5 6 7 8 9 10



1 2 3 4 5 6 7 8 9 10

Fig. 2

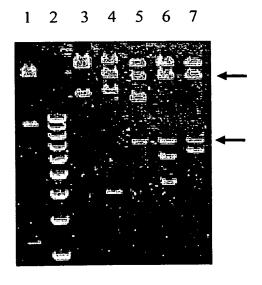


Fig. 3

Fig. A

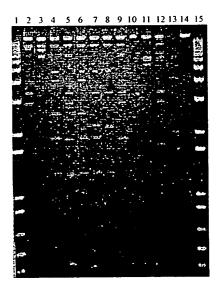


Fig. B



Fig. 4

GAGCTCTGAA	CCGTGGAAAC	GAACATGACC	CTTGCCTGCC	TGCTTCCCTG	GGTGGGTCAA	GGGTAATGAA	70
GTGGTGTGCA	GGAAATGGCC	ATGTAAATTA	CACGACTCTG	CTGATGGGGA	CCGTTCCTTC	CATCATTATT	140
CATCTTCACC	CCCAAGGACT	GAATGATTCC	AGCAACTTCT	TCGGGTGTGA	CAAGCCATGA	CAAAACTCAG	210
TACAAACACC	ACTCTTTTAC	TAGGCCCACA	GAGCACGGGC	CACACCCCTG	ATATATTAAG	AGTCCAGGAG	280
AGATGAGGCT	GCTTTCAGCC	ACCAGGCTGG	GGTGACAACA	GCGGCTGAAC	AGTCTGTTCC	TCTAGACTAG	350
TAGACCCTGG	CAGGCACTCC	CCCAAATTCT	AGGGCCTGGT	TGCTGCTTCC	CGAGGGCGCC	ATCTGCCCTG	420
GAGACTCAGC	CTGGGGTGCC	ACACTGAGGC	CAGCCCTGTC	TCCACACCCT	CCGCCTCCAG	GCCTCAGCTT	490
CTCCAGCAGC	TTCCTAAACC	CTGGGTGGGC	CGTGTTCCAG	CGCTACTGTC	TCACCTGTCC	CACTGTGTCT	560
TGTCTCAGCG	ACGTAGCTCG	CACGGTTCCT	CCTCACATGG	GGTGTCTGTC	TCCTTCCCCA	ACACTCACAT	630
GCGTTGAAGG	GAGGAGATTC	TGCGCCTCCC	AGACTGGCTC	CTCTGAGCCT	GAACCTGGCT	CGTGGCCCCC	700
GATGCAGGTT	CCTGGCGTCC	GGCTGCACGC	TGACCTCCAT	TTCCAGGCGC	TCCCCGTCTC	ĊTGTCATCTG	770
CCGGGGCCTG	CCGGTGTGTT	CTTCTGTTTC	TGTGCTCCTT	TCCACGTCCA	GCTGCGTGTG	TCTCTGCCCG	840
CTAGGGTCTC	GGGGTTTTTA	TAGGCATAGG	ACGGGGGCGT	GGTGGGCCAG	GGCGCTCTTG	GGAAATGCAA	910
CATTTGGGTG	TGAAAGTAGG	AGTGCCTGTC	CTCACCTAGG	TCCACGGGCA	CAGGCCTGGG	GATGGAGCCC	980
CCGCCAGGGA	CCCGCCCTTC	TCTGCCCAGC	ACTTTCCTGC	CCCCTCCCT	CTGGAACACA	GAGTGGCAGT	1050
TTCCACAAGC	ACTAAGCATC	CTCTTCCCAA	AAGACCCAGC	ATTGGCACCC	CTGGACATTT	GCCCCACAGC	1120
CCTGGGAATT	CACGTGACTA	CGCACATCAT	GTACACACTC	CCGTCCACGA	CCGACCCCCG	CTGTTTTATT	1190
TTAATAGCTA	CAAAGCAGGG	AAATCCCTGC	TAAAATGTCC	TTTAACAAAC	TGGTTAAACA	AACGGGTCCA	1260
TCCGCACGGT	GGACAGTTCC	TCACAGTGAA	GAGGAACATG	CCGTTTATAA	AGCCTGCAGG	CATCTCAAGG	1330
GAATTACGCT	GAGTCAAAAC	TGCCACCTCC	ATGGGATACG	TACGCAACAT	GCTCAAAAAG	AAAGAATTTC	1400
ACCCCATGGC	AGGGGAGTGG	TTAGGGGGGT	TAAGGACGGT	GGGGGCGCA	GCTGGGGGCT	ACTGCACGCA	1470
CCTTTTACTA	AAGCCAGTTT	CCTGGTTCTG	ATGGTATTGG	CTCAGTTATG	GGAGACTAAC	CATAGGGGAG	1540
TGGGGATGGG	GGAACCCGGA	GGCTGTGCCA	TCTTTGCCAT	GCCCGAGTGT	CCTGGGCAGG	ATAATGCTCT	1610
AGAGATGCCC	ACGTCCTGAT	TCCCCCAAAC	CTGTGGACAG	AACCCGCCCG	GCCCCAGGGC	CTTTGCAGGT	1680
GTGATCTCCG	TGAGGACCCT	GAGGTCTGGG	ATCCTTCGGG	ACTACCTGCA	GGCCCGAAAA	GTAATCCAGG	1750
GGTTCTGGGA	AGAGGCGGC	AGGAGGGTCA	GAGGGGGCA	GCCTCAGGAC	GATGGAGGCA	GTCAGTCTGA	1820
GGCTGAAAAG	GGAGGGAGGG	CCTCGAGCCC	AGGCCTGCAA	GCGCCTCCAG	AAGCTGGAAA	AAGCGGGGAA	1890
GGGACCCTCC	ACGGAGCCTG	CAGCAGGAAG	GCACGGCTGG	CCCTTAGCCC	ACCAGGGCCC	ATCGTGGACC	1960
TCCGGCCTCC	GTGCCATAGG	AGGGCACTCG	CGCTGCCCTT	CTAGCATGAA	GTGTGTGGGG	ATTTGCAGAA	2030
GCAACAGGAA	ACCCATGCAC	TGTGAATCTA	GGATTATTTC	AAAACAAAGG	TTTACAGAAA	CATCCAAGGA	2100
CAGGGCTGAA	GTGCCTCCGG	GCAAGGCAG	GGCAGGCACG	AGTGATTTTA	TTTAGCTATT	TTATTTTATT	2170
TACTTACTTT	CTGAGACAGA	GTTATGCTCT	TGTTGCCCAG	GCTGGAGTGC	AGCGGCATGA	TCTTGGCTCA	2240
CTGCAACCTC	CGTCTCCTGG	GTTCAAGCAA	TTCTCGTGCC	TCAGCCTCCC	AAGTAGCTGG	GATTTCAGGC	2310
GTGCACCACC	ACACCCGGCT	AATTTTGTAT	TTTTAGTAGA	GATGGGCTTT	CACCATGTTG	GTCAAGCTGA	2380
TCTCAAAATC	CTGACCTCAG	GTGATCCGCC	CACCTCAGCC	TCCCAAAGTG	CTGGGATTAC	AGGCATGAGC	2450
CACTGCACCT	GGCCTATTTA	ACCATTTTAA	AACTTCCCTG	GGCTCAAGTC	ACACCCACTG	GTAAGGAGTT	2520
CATGGAGTTC	AATTTCCCCT	TTACTCAGGA	GTTACCCTCC	TTTGATATTT	TCTGTAATTC	TTCGTAGACT	2590
GGGGATACAC	CGTCTCTTGA	CATATTCACA	GTTTCTGTGA	CCACCTGTTA	TCCCATGGGA	CCCACTGCAG	2660
GGGCAGCTGG	GAGGCTGCAG	GCTTCAGGTC	CCAGTGGGGT	TGCCATCTGC	CAGTAGAAAC	CTGATGTAGA	2730
ATCAGGGCGC	AAGTGTGGAC	ACTGTCCTGA	ATCTCAATGT	CTCAGTGTGT	GCTGAAACAT	GTAGAAATTA	2800
AAGTCCATCC	CTCCTACTCT	ACTGGGATTG	AGCCCCTTCC	CTATCCCCC	CCAGGGGCAG	AGGAGTTCCT	2870
CTCACTCCTG	TGGAGGAAGG	AATGATACTT	TGTTATTTTT	CACTGCTGGT	ACTGAATCCA	CTGTTTCATT	2940
TGTTGGTTTG	TTTGTTTTGT	TTTGAGAGGC	GGTTTCACTC	TTGTTGCTCA	GGCTGGAGGG	AGTGCAATGG	3010
CGCGATCTTG	GCTTACTGCA	GCCTCTGCCT	CCCAGGTTCA	AGTGATTCTC	CTGCTTCCGC	CTCCCATTTG	3080
GCTGGGATTA	CAGGCACCCG	CCACCATGCC	CAGCTAATTT	TTTGTATTTT	TAGTAGAGAC	GGGGGTGGGT	3150

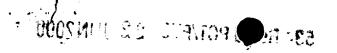


Fig. 4 (continuation)

GGGGTTCACC	ATGTTGGCCA	GGCTGGTCTC	GAACTTCTGA	CCTCAGATGA	TCCACCTGCC	TCTGCCTCCT	3220
AAAGTGCTGG	GATTACAGGT	GTGAGCCACC	ATGCCCAGCT	CAGAATTTAC	TCTGTTTAGA	AACATCTGGG	3290
TCTGAGGTAG	GAAGCTCACC	CCACTCAAGT	GTTGTGGTGT	TTTAAGCCAA	TGATAGAATT	TTTTTATTGT	3360
TGTTAGAACA	CTCTTGATGT	TTTACACTGT	GATGACTAAG	ACATCATCAG	CTTTTCAAAG	ACACACTAAC	3430
TGCACCCATA	ATACTGGGGT	GTCTTCTGGG	TATCAGCAAT	CTTCATTGAA	TGCCGGGAGG	CGTTTCCTCG	3500
CCATGCACAT	GGTGTTAATT	ACTCCAGCAT	AATCTTCTGC	TTCCATTTCT	TCTCTTCCCT	CTTTTAAAAT	3570
TGTGTTTTCT	ATGTTGGCTT	CTCTGCAGAG	AACCAGTGTA	AGCTACAACT	TAACTTTTGT	TGGAACAAAT	3640
TTTCCAAACC	GCCCCTTTGC	CCTAGTGGCA	GAGACAATTC	ACAAACACAG	CCCTTTAAAA	AGGCTTAGGG	3710
ATCACTAAGG	GGATTTCTAG	AAGAGCGACC	TGTAATCCTA	AGTATTTACA	AGACGAGGCT	AACCTCCAGC	3780
GAGCGTGACA	GCCCAGGGAG	GGTGCGAGGC	CTGTTCAAAT	GCTAGCTCCA	TAAATAAAGC	AATTTCCTCC	3850
GGCAGTTTCT	GAAAGTAGGA	AAGGTTACAT	TTAAGGTTGC	GTTTGTTAGC	ATTTCAGTGT	TTGCCGACCT	3920
CAGCTACAGC	ATCCCTGCAA	GGCCTCGGGA	GACCCAGAAG	TTTCTCGCCC	CCTTAGATCC	AAACTTGAGC	3990
AACCCGGAGT	CTGGATTCCT	GGGAAGTCCT	CAGCTGTCCT	GCGGTTGTGC	CGGGGCCCCA	GGTCTGGAGG	4060
GGACCAGTGG	CCGTGTGGCT	TCTACTGCTG	GGCTGGAAGT	CGGGCCTCCT	AGCTCTGCAG	TCCGAGGCTT	4130
GGAGCCAGGT	GCCTGGACCC	CGAGGCTGCC	CTCCACCCTG	TGCGGGCGGG	ATGTGACCAG	ATGTTGGCCT	4200
CATCTGCCAG	ACAGAGTGCC	GGGGCCCAGG	GTCAAGGCCG	TTGTGGCTGG	TGTGAGGCGC	CCGGTGCGCG	4270
GCCAGCAGGA	GCGCCTGGCT	CCATTTCCCA	CCCTTTCTCG	ACGGGACCGC	CCCGGTGGGT	GATTAACAGA	4 3 4.0
TTTGGGGTGG	TTTGCTCATG	GTGGGGACCC	CTCGCCGCCT	GAGAACC'IGC	AAAGAGAAAT	GACGGGCCTG	4410
TGTCAAGGAG	CCCAAGTCGC	GGGGAAGTGT	TGCAGGGAGG	CACTCCGGGA	GGTCCCGCGT	GCCCGTCCAG	4480
GGAGCAATGC	GTCCTCGGGT	TCGTCCCCAG	CCGCGTCTAC	GCGCCTCCGT	CCTCCCCTTC	ACGTCCGGCA	4550
TTCGTGGTGC	CCGGAGCCCG	ACGCCCCGCG	TCCGGACCTG	GAGGCAGCCC	TGGGTCTCCG	GATCAGGCCA	4620
GCGGCCAAAG	GGTCGCCGCA	CGCACCTGTT	CCCAGGGCCT	CCACATCATG	GCCCCTCCCT	CGGGTTACCC	4590
CACAGCCTAG	GCCGATTCGA	CCTCTCTCCG	CTGGGGCCCT	CGCTGGCGTC	CCTGCACCCT	GGGAGCGCGA	4760
GCGGCGCGCG	GGCGGGGAAG	CGCGGCCCAG	ACCCCCGGGT	CCGCCCGGAG	CAGCTGCGCT	GTCGGGGCCA	4830
GGCCGGGCTC	CCAGTGGATT	CGCGGGCACA	GACGCCCAGG	ACCGCGCTCC	CCACGTGGCG	GAGGGACTGG	4900
GGACCCGGGC	ACCCGTCCTG	CCCCTTCACC	TTCCAGCTCC	GCCTCCTCCG	CGCGGACCCC	GCCCGTCCC	4970
GACCCCTCCC	GGGTCCCCGG	CCCAGCCCCC	TCCGGGCCCT	CCCAGCCCCT	CCCCTTCCTT	TCCGCGGCCC	5040
CGCCCTCTCC	TCGCGGCGCG	AGTTTCAGGC	AGCGCTGCGT	CCTGCTGCGC	ACGTGGGAAG	CCCTGGCCCC	6110
GGCCACCCC	GCG <u>ATG</u>						6126

Fig. 5

1 G ni ingres See value